Current Conditions Biota Sampling 2019

April 4, 2019

DQOs for Biota Tissue Program

Primary DQO – Evaluate trends

Collect initial data to establish current chemical (TCDD and PCBs) concentrations in fish and crab in the LPRSA upper 9-mile area (RM 8.3 to Dundee Dam) to monitor changes in post-interim action changes in tissue concentrations.

Secondary DQO – Refine/validate FWM

Evaluate FWM performance using the 2019 fish and crab tissue data and refine model as needed.

Current Conditions DQOs: Fish and crab tissue analysis

| DQO | Study Questions | Data Evaluations |
|-----------------------------------|--|---|
| 1 – Evaluate trends | What are the current chemical (TCDD, PCBs) concentrations in fish and crab in the LPRSA (upper 9 miles)? How do the current conditions determine potential for biota recovery via trend analysis? | Establish concentrations for future trend analysis |
| 2 – Refine and validate FWM | Hoes does the FWM perform using the 2019 data? Can the calibration of the model be improved? | Evaluate model performanceRefine FWM |

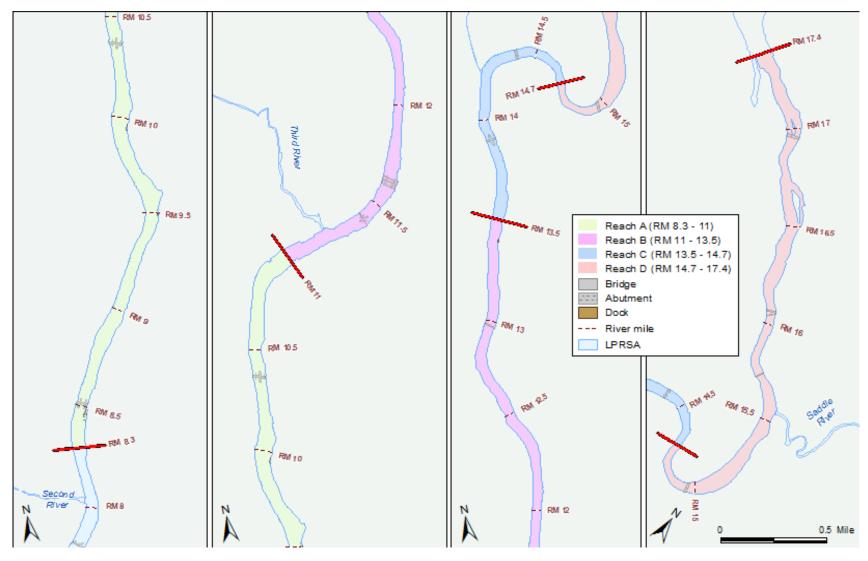
Overview of Approach

- Objective understand area-wide average TCDD and total PCB concentrations in fish and crab for the upper 9 miles of the LPRSA (RM 8.3-Dam).
- Sampling areas
 - Focus on four primary areas for sample design informed by 2009/2010 data and Food Web Model (FWM) areas
 - RM 8.3-11, RM 11-13.5, RM 13.5-14.7, and RM 14.7-Dam
- Collect samples during the same time of year as most samples in 2009/2010 (late summer/early fall).

Selection of Species for Sampling

| Possible Species | Useful for FWM Validation? | LPRSA Abundance | Site Fidelity (short-term) | Salinity Tolerance | Proposed for Sampling? |
|-----------------------------|----------------------------|--------------------------------|--|-----------------------|------------------------|
| Small forage fish (sunfish) | X | moderate/ high | high | no | Yes |
| American eel (<35 cm) | X | moderate (high near dam) | moderate; move out of LPRSA to spawn as adults | yes | No |
| American eel (>35 cm) | Х | moderate | moderate; move out of LPRSA to spawn as adults | yes | Yes |
| Blue crab | X | high | low/moderate | yes | Yes |
| Carp | X | high | moderate; may also move into tributaries | no | Yes |
| White perch (adult) | Х | moderate | low | yes | yes |
| Catfish | X | moderate | low | moderate | No |
| Bass | X | low | high | no | No |

Sampling Areas



Evaluating Trends

- Methods:
 - Graphical evaluation
- Reducing variance
 - Target fish sizes
 - Species
 - Sampling areas

Proposed biota tissue sampling

Proposed species for sampling:

| Species | Proposed Tissue Type(s) | Target No. of Samples | Fish Per Composite | Most Effective Sampling Method (s) |
|-----------------------------|------------------------------------|--------------------------|-----------------------|--|
| American eel (35-60 cm) | Fillet, remainder (calc'd WB) | 12 composites | 3 | trotline |
| Blue crab | muscle/hep, carcass (calc'd WB) | 12 composites | 3 | crab trap, gillnet |
| Carp | Fillet, remainder (calc'd WB) | 12 composites | 3 | gillnet, boat electrofishing |
| Small forage fish (sunfish) | Whole body | 12 composites | 5 to 10 | electrofishing (boat/backpack), minnow trap, beach seine |
| Perch | Fillet, remainder (calc'd WB) | 12 composites | 3 | gillnet, boat electrofishing |

- Year 1 target maximum of 60 composites (108 analytical samples)
- Methods trotlines, boat electrofishing, crab traps, minnow traps, gillnets, and beach seine

Adaptive Sample Design

Use Year 1 data to determine the following:

Tissue types –

• Is the relationship between fillet (or muscle for blue crab) and whole body concentrations sufficient to allow for only the collection of fillet (or muscle) data during year 3 sampling?

Need for Year 2 Sampling and Number of samples –

- Evaluate the need for Year 2 sampling (i.e., conduct sampling if significant changes in conditions that would affect chemical exposure such as shift in water temperatures or sustained water flow that might affect prey availability.)
- Are fewer samples acceptable for future monitoring work based on measured variances in tissue concentrations?

Biota tissue sampling task deadlines

| Month | Task |
|-----------|--|
| | Determine details of sampling plan with EPA |
| | Finalize boat/equipment contractor(s) |
| April | Finalize laboratories and analytical methods |
| | Finalize field sampling dates |
| | Initiate work on parts of QAPP Addendum |
| | QAPP Addendum Preparation and submittal to CPG |
| May | Establish subcontract agreements and POs (boat operators, laboratories, and validator) |
| | Apply for NJDEP scientific collection permit |
| June | EPA review of QAPP Addendum |
| Luter | Locate and set-up field laboratory. |
| July | Conduct pre-sampling recon (site conditions and access) |
| August | Mobilization of supplies and equipment prep to field facility |
| August | Finalize QAPP addendum |
| September | Fieldwork begins (target starting field effort in mid-September) |

QAPP Addendum Schedule

• Draft schedule for QAPP Addendum:

| | QAPP Addendum | Review Duration |
|------------------------------|---------------|-----------------|
| To CPG | May 17 | 2 weeks |
| To EPA | June 15 | 3 weeks |
| Revised to EPA (draft final) | July 15 | 1 week |
| Comments from EPA | August 1 | 1 week |
| Final QAPP to EPA | August 21 | |

 Assumes meetings with EPA prior to May 1 to finalize key details (e.g., species, number of samples, laboratories/methods, etc.)

Analytical Laboratories

| Analysis/ Sample Prep | Previous Lab | Proposed Lab |
|-----------------------------------|-------------------------------|-------------------------------|
| Tissue processing and compositing | Alpha Analytical | Alpha Analytical |
| PCBs | SGS - Analytical Perspectives | Cape Fear Analytical |
| Dioxins/furans | SGS - Analytical Perspectives | Cape Fear Analytical |
| Lipids | CAS | TBD – Method dependent |
| Percent Moisture | Alpha Analytical | Cape Fear or Alpha Analytical |

Notes:

- Is Bligh-Dyer lipid method required?
- Staffing changes at SGS-AP
- Validator = LDC
- Data management = ddms